

REPLACEMENT
SHEET

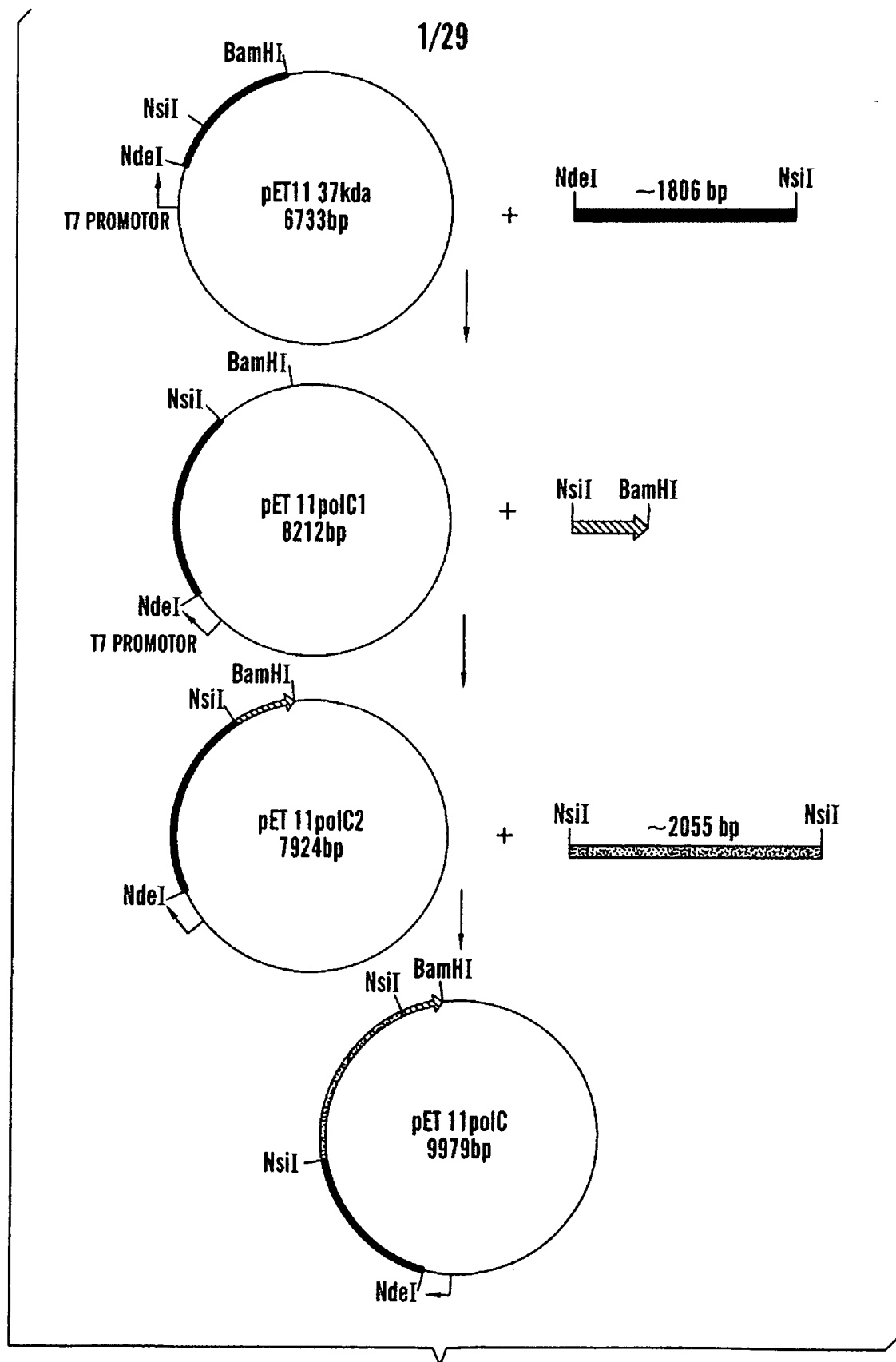


FIG. 1

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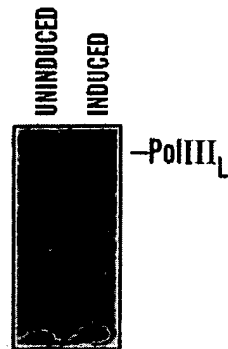


FIG. 2A

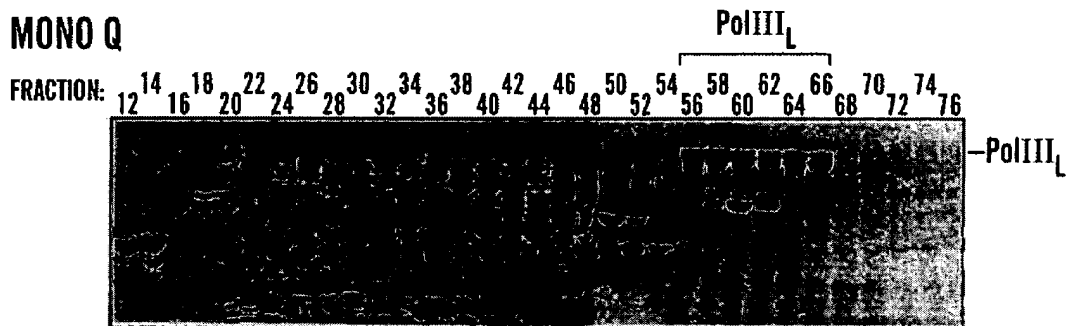


FIG. 2B

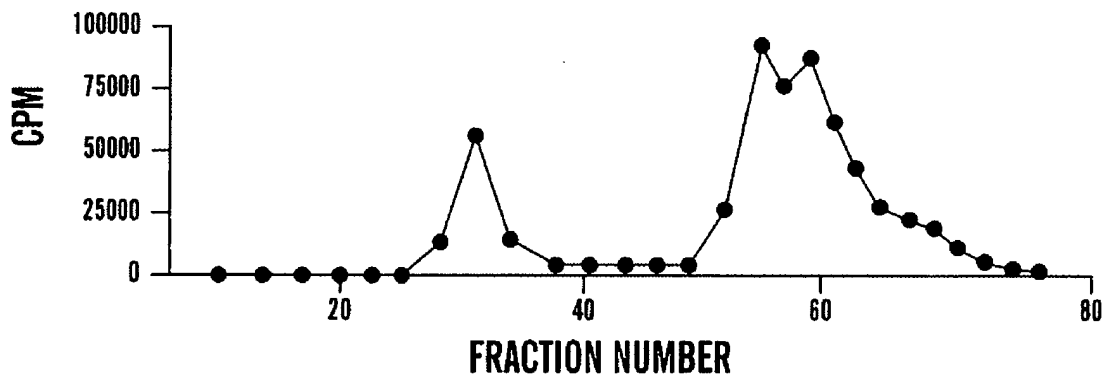


FIG. 2C

PHOSPHOCELLULOSE

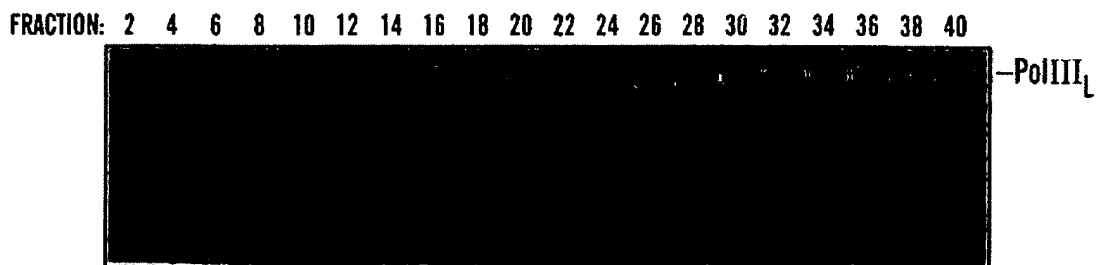


FIG. 2D

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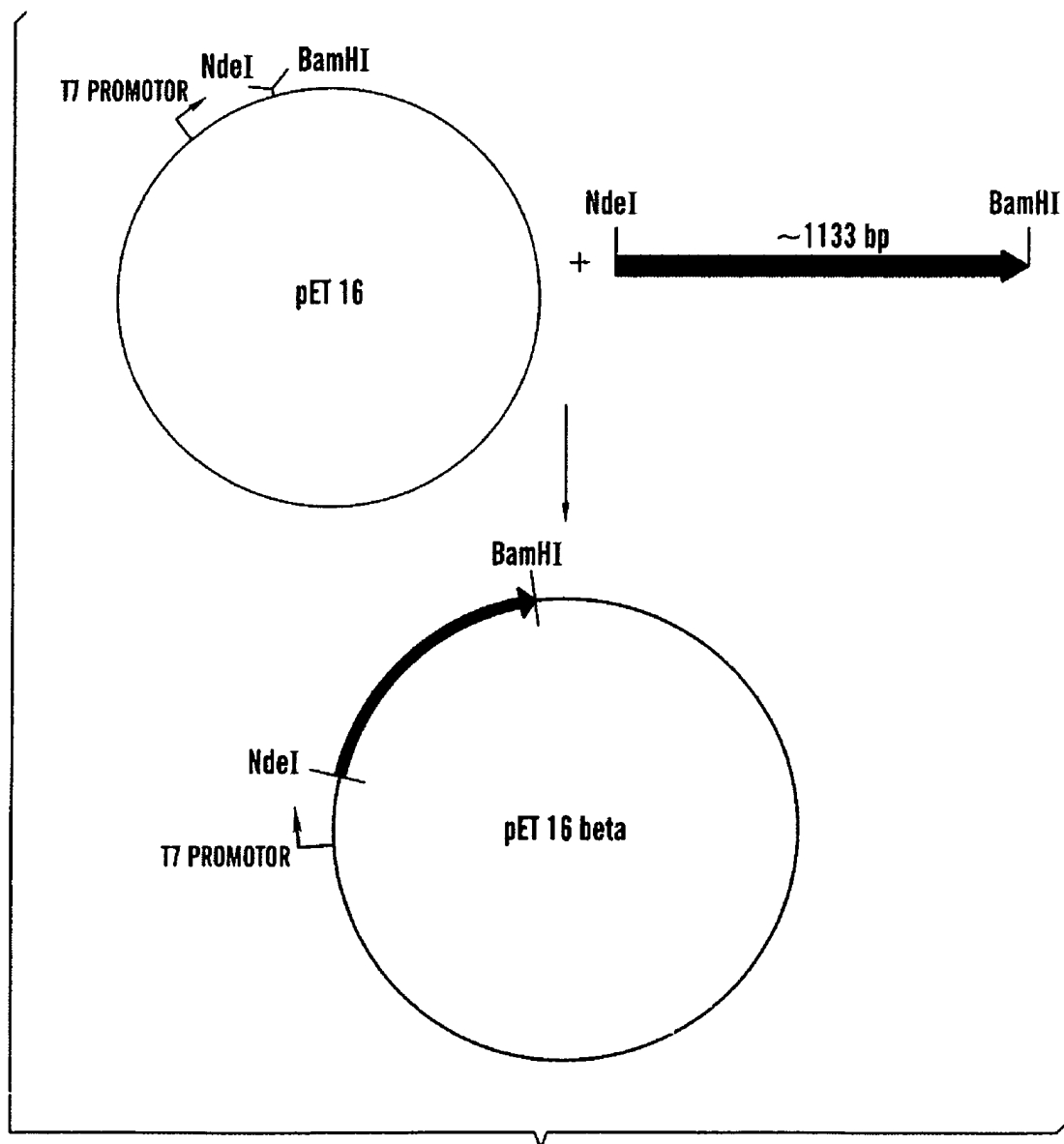


FIG. 3

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INDUCTION

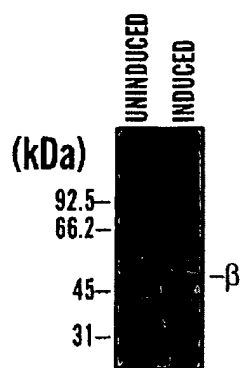


FIG. 4A

NICKEL COLUMN

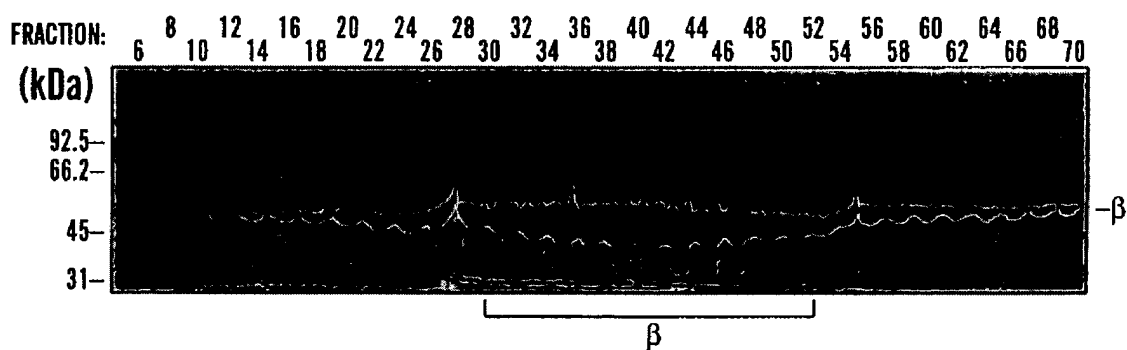


FIG. 4B

MONO Q

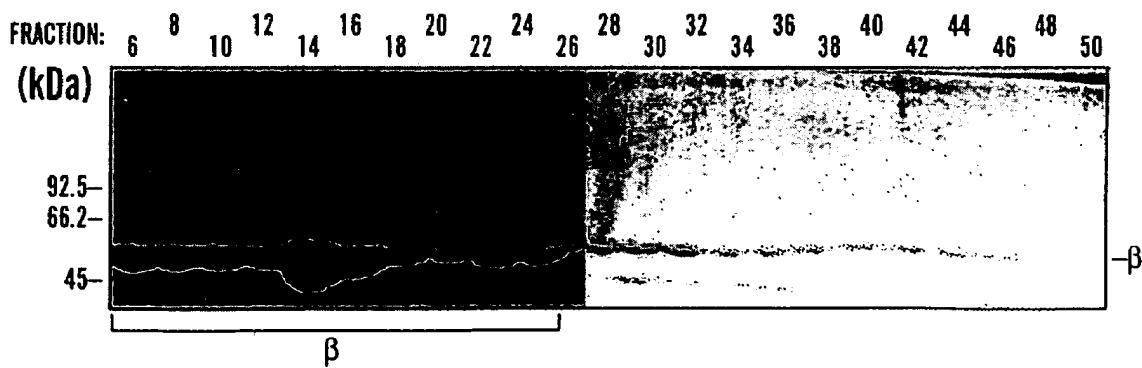


FIG. 4C

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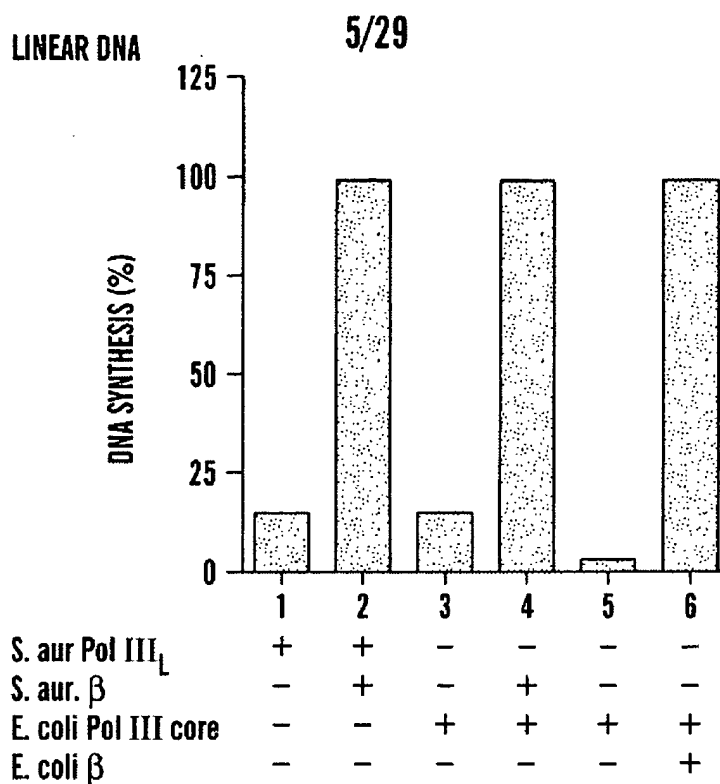


FIG. 5A

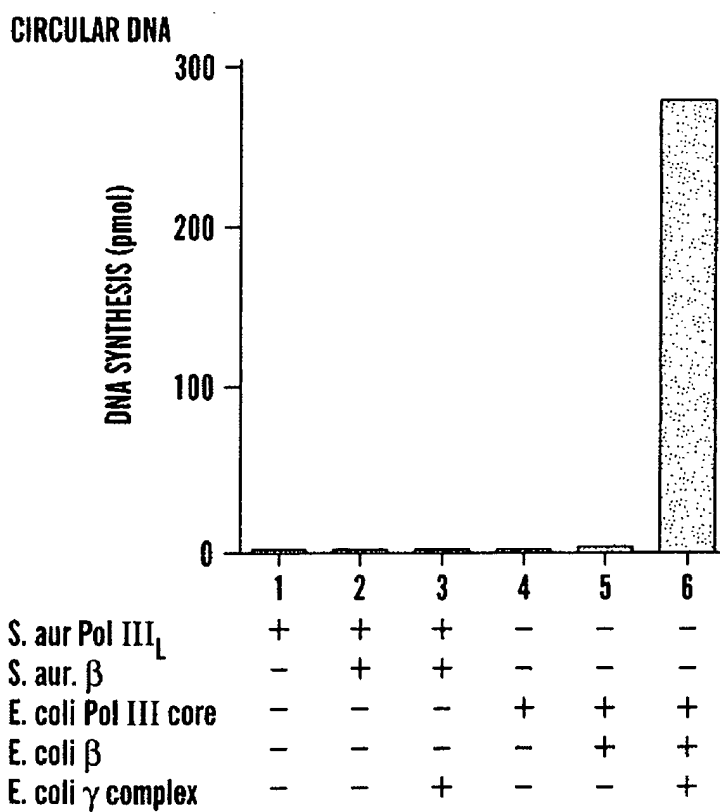


FIG. 5B

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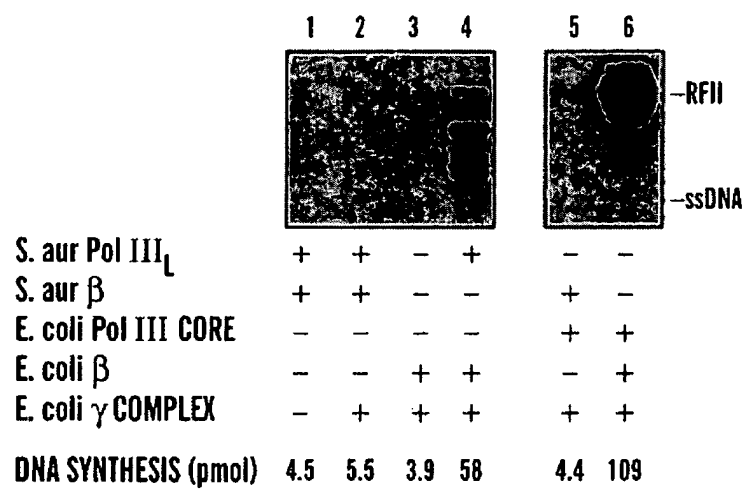


FIG. 6

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MonoQ

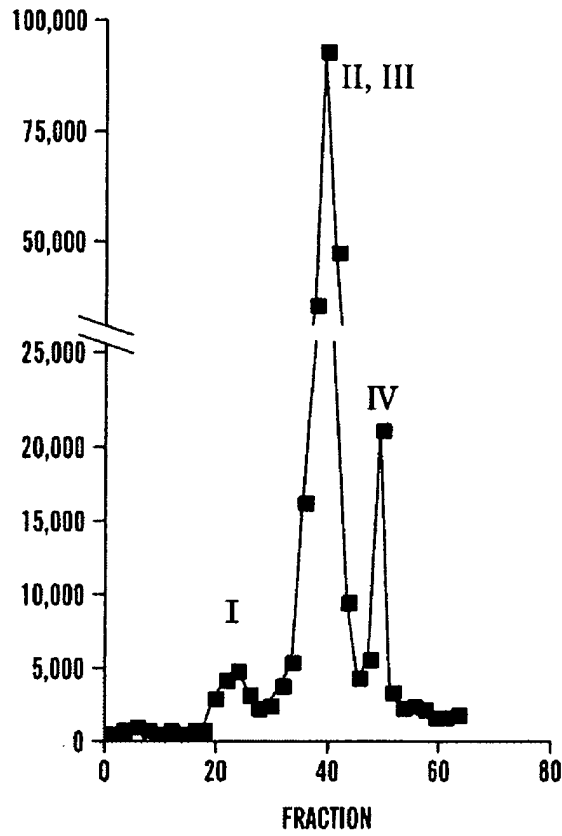


FIG. 7A

P-Cell

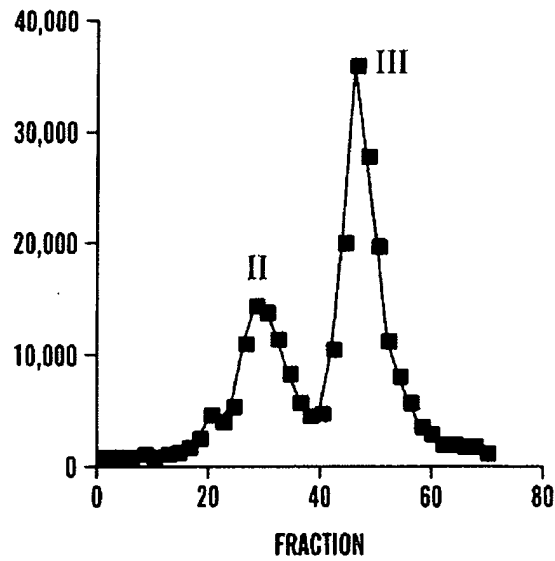


FIG. 7B

REPLACEMENT SHEET

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AGAROSE GEL

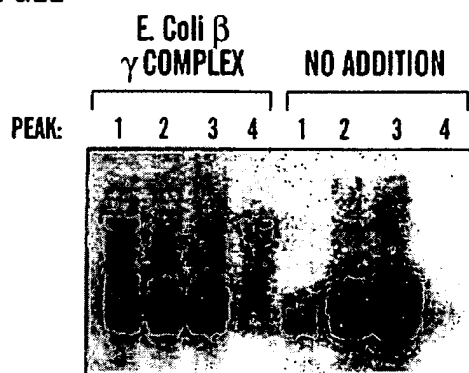


FIG. 8A

DNA SYNTHESIS

ADDITION	DNA SYNTHESIS (PMOI) PEAK			
	PEAK 1	PEAK 2	PEAK 3	PEAK 4
NONE	22.7	70.6	146.1	4.7
E. coli β , γ COMPLEX	72.9	61.2	71.4	25.9

FIG. 8B

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S. aureus	KIWRATCIWNCDFRSSACKAVAKDVGRIMGFDEVTLNEISSLI PHKLGITLDEAYQID-D
E. coli	MYGRDAVSQIIITFGTMAAKAVIRDVGRVLGHPYGFVDRI SKLIPDPGMTLAKAFEAPQ
Sal. typ	MYGRDAVSQIIITFGTMAAKAVIRDVGRVLGHPYGFVDRI SKLVPDPGMTLAKAFEAPQ
S. aureus	FKKFVHRNHRHQRWFSICKKLEGLPRHTSTHAAGI IINDHPLYEYAPLTCKGDTG--LLTQ
E. coli	LPEIYEADEEVKALIDMARKLEGVTRNAGKHAGGVIAPTKITDFAPLYCDEEGKHPVTQ
Sal. typ	LPEIYEADEEVRALIDMARKLEGVTRNAGKHAGGVIAPTKITDFAPLYCDEEGKHPVTQ
S. aureus	WMTPEAERIGLLKIDFLGLRNL SIHQILTRVEKDLGN-----IDIEKIPFDDQKVFELI
E. coli	FDKSDVEYAGLVKFDLGLR TLTIINWALEMINKRRRAKNGEPPLDIAAIPLDDKKSFDML
Sal. typ	FDKSDVEYAGLVKFDLGLR TLTIINWALEMINKRRRAKNGEPPLDIAAIPLDDKKSFDML
S. aureus	SQGDTTGIFQLES DGVSVLKCLKPEHFEDIVAVTSLYRPGPMEE--IPTYITRRHDP S-
E. coli	QRSETTAVFQLESRGMDLIKRLQPCDFEDMIALVALFRPGPLQSGMVDNFIDRKHGREE
Sal. typ	QRSETTAVFQLESRGMDLIKRLQPCDFEDMIALVALFRPGPLQSGMVDNFIDRKHGREE
S. aureus	-----KVQYLHPHLEPILKN TYGVITYQEQIMQIASTFANFSYGEADILRRAMSKKNRAVL
E. coli	ISYPDVQWQHESLKPVLEPTYGIILYQEQVMQIAQVLSGYTLGGADMLRRAMGKKKPEEM
Sal. typ	LSYPDVQWQHESLKPVLEPTYGIILYQEQVMQIAQVLSGYTLGGADMLRRAMGKKKPEEM
S. aureus	ERDAQHFIEG TKQNGYHEDI SKQIFDLI-----
E. coli	AKQRSVFAEGA EKNGINAE LAMKIFDLVEKFPAGYGFNKSHSAAYALVSYQTLWLKAHYPA
Sal. typ	AKQRSVFEEGAKNGIDGELAMKIFDLVEKFPAGYGFNKSHSAAYALVSYQTLWLKAHYPA

FIG. 9

REPLACEMENT SHEET

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ATP site

S. aureus
 B. sub.
 E. coli

MKGYCLWRCNLDYQALFVVTP--KFEDVVGQEHSEDCAMG-----SHAYLFSGPRGTGKT
 -----MSYQALYRVFRPQRFEDVVGQEHITKTLQNALLOKKFSHAYLFSGPRGTGKT
 -----MSYQVLARKWRPQTFADVVGQEHVLTALANGLSLGRIHHAYLFSGTRGVGKT
 ** * * * * *
 * * * * *
 * * * * *

Zn++ finger

S. aureus
 B. sub
 E. coli

SIAKVFAKAINCLNSTDGEPCNECHICKGITQGTNSDVIEIDAASNNGVDEIRNIRDVKVYA
 SAAKIFAKAVNCEHAPVDEPCNECAACKGITNGSISDVIEIDAASNNGVDEIRDIRDKVKFA
 SIARLLAKGLNCETGITATPCGVCDNCREIEQGRFVDLIEIDAASRTKVEDTRDLLDNVQYA
 * * * * *
 * * * * *
 * * * * *

S. aureus
 B. sub
 E. coli

PSESKYKVYIIDEVHMLTTGAFNALLKTLLEPPPAHAIFILATTEPHKIPPTIISRA
 PSAVTYKVYIIDEVHMLSIGAFNALLKTLLEPPPEHCIFILATTEPHKIPPTIISRC
 PARGRFKVYIIDEVHMLSRHSFNALLKTLLEPPPEHVKFLLATDPQKLPVTLISRC
 * * * * *
 * * * * *
 * * * * *

FIG. 10

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FIG. 11

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FIG. 12A

FIG. 12B

REPLACEMENT SHEET

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<i>S. aureus</i> α -L/ β :	+	+	+
<i>S. aureus</i> $\tau\delta\delta'$ (ng):	-	30	90

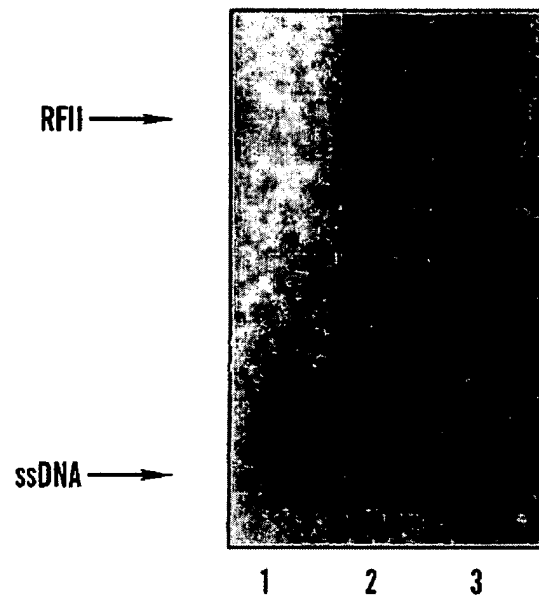


FIG. 13

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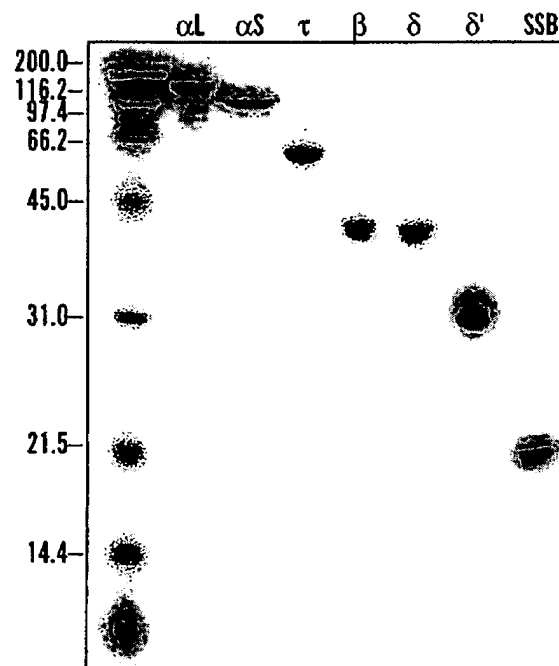


FIG. 14

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SUPEROSE 6

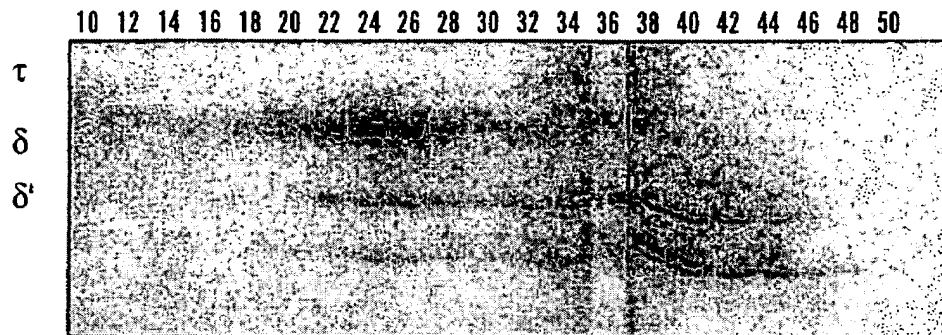


FIG. 15A

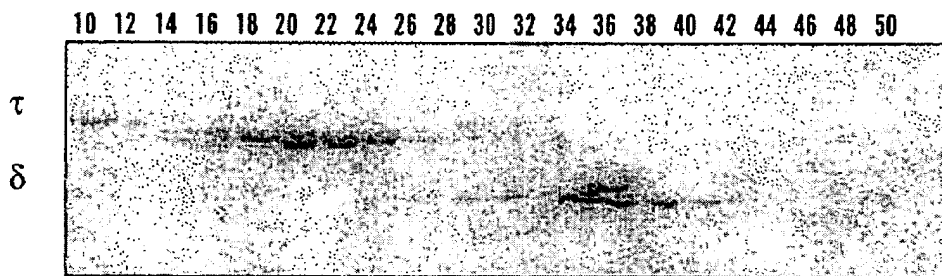


FIG. 15B

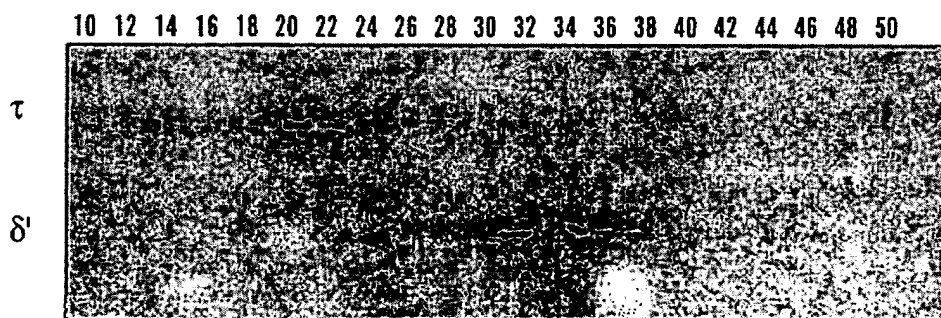


FIG. 15C

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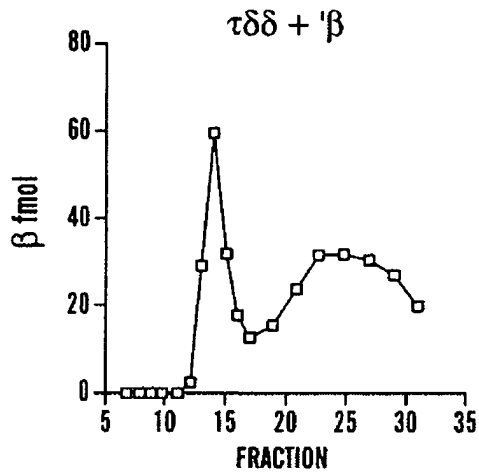


FIG. 16A

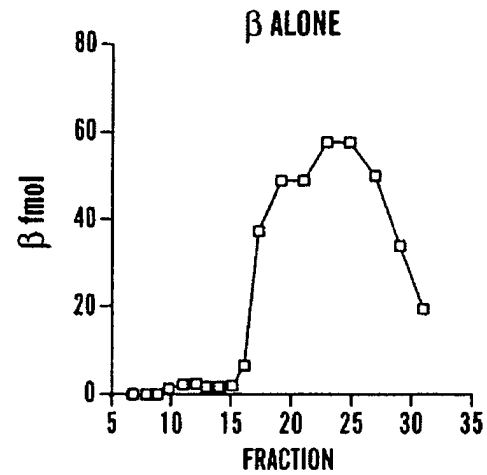


FIG. 16B

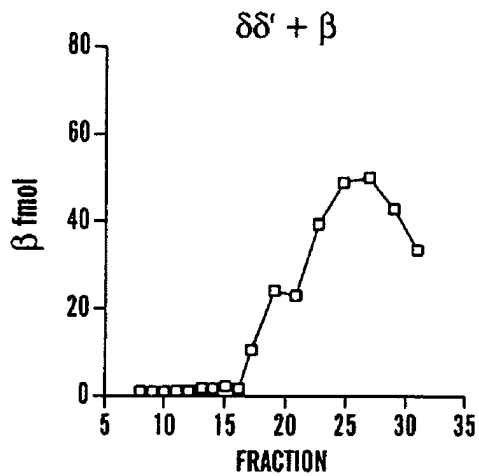


FIG. 16C

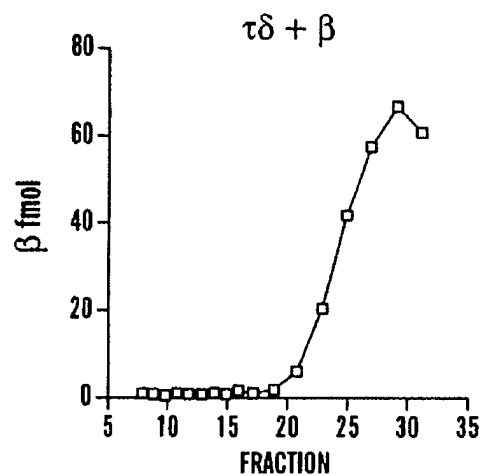


FIG. 16D

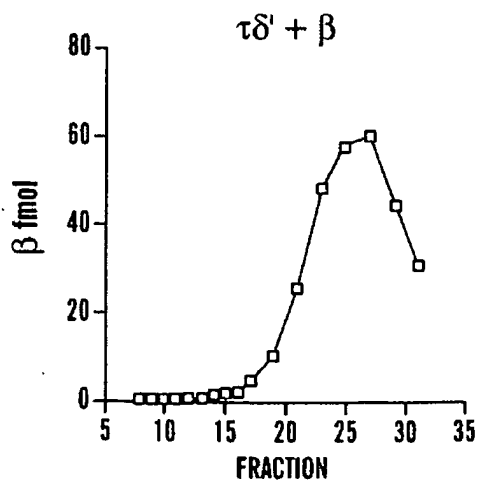
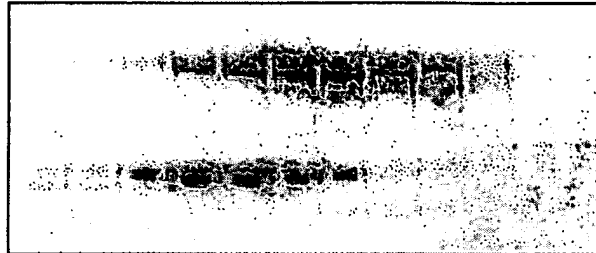


FIG. 16E

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24 26 28 30 32 34 36 38 40 42 44 46 48

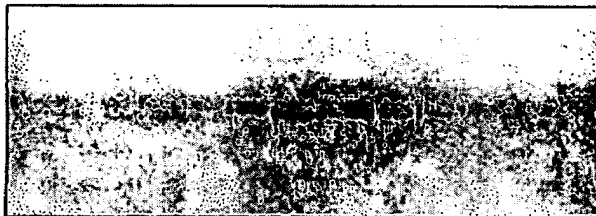


← α LARGE

← τ

FIG. 17A

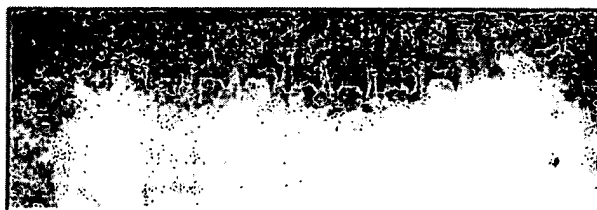
24 26 28 30 32 34 36 38 40 42 44 46 48



← τ

FIG. 17B

24 26 28 30 32 34 36 38 40 42 44 46 48



← α LARGE

FIG. 17C

REPLACEMENT
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SUPEROSE 6

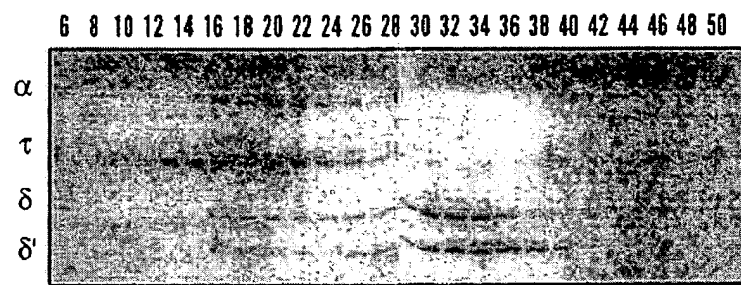


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FIG. 19

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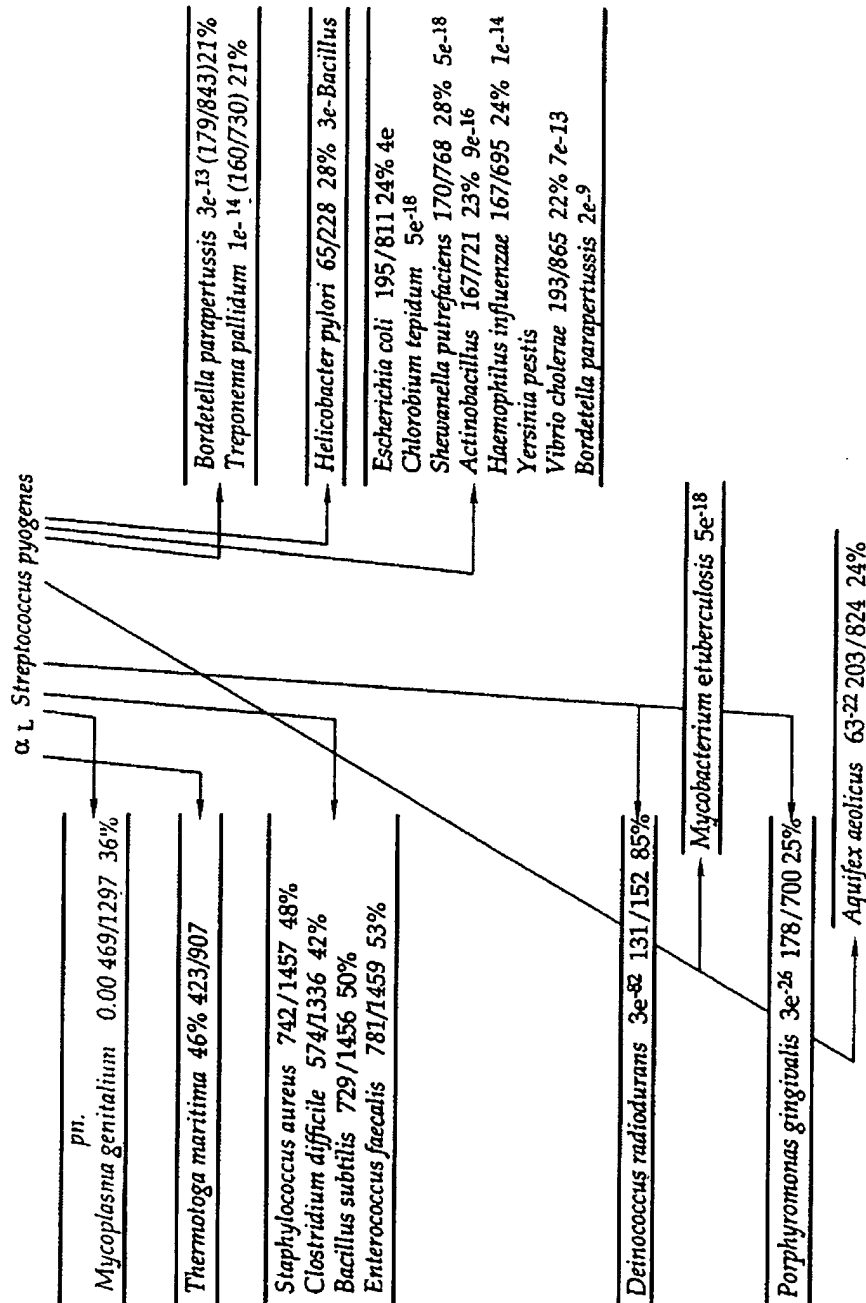


FIG. 20A

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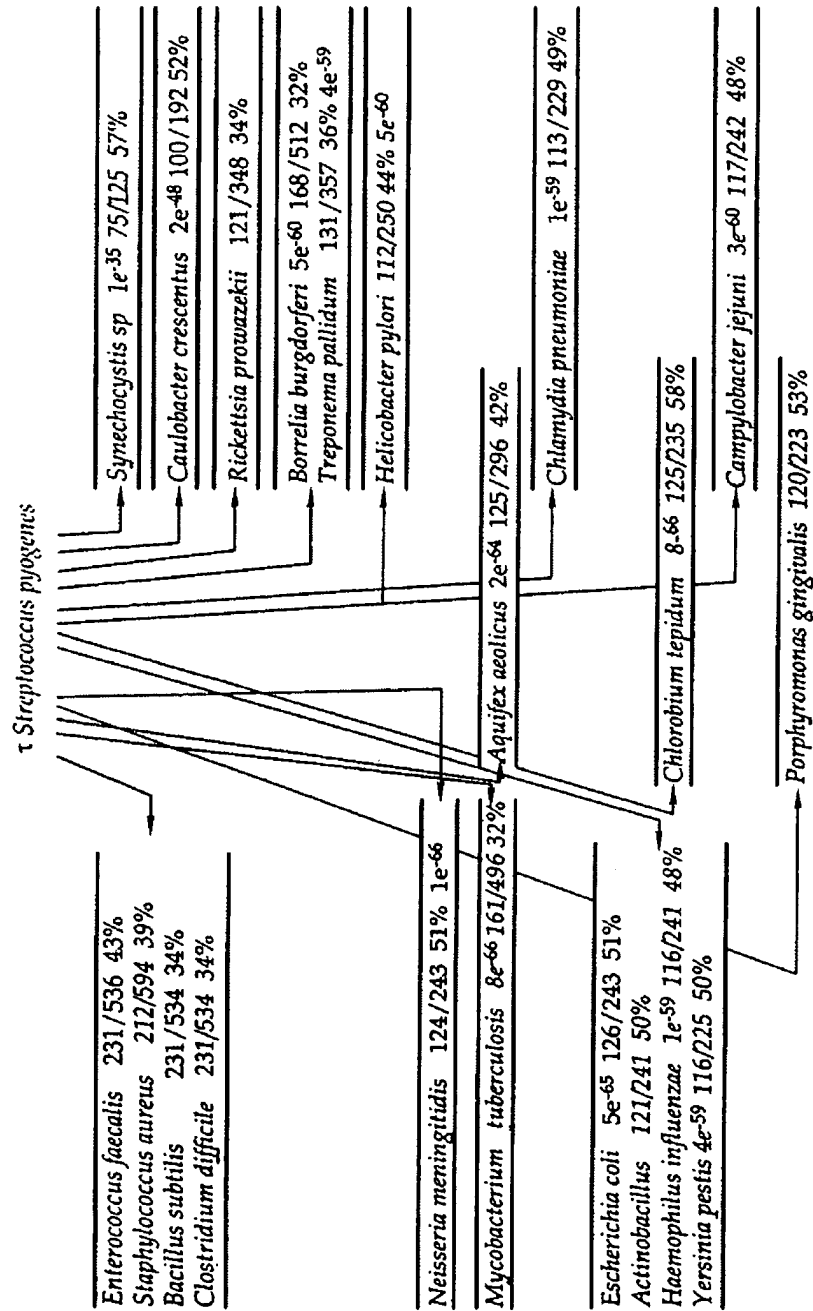


FIG. 20B

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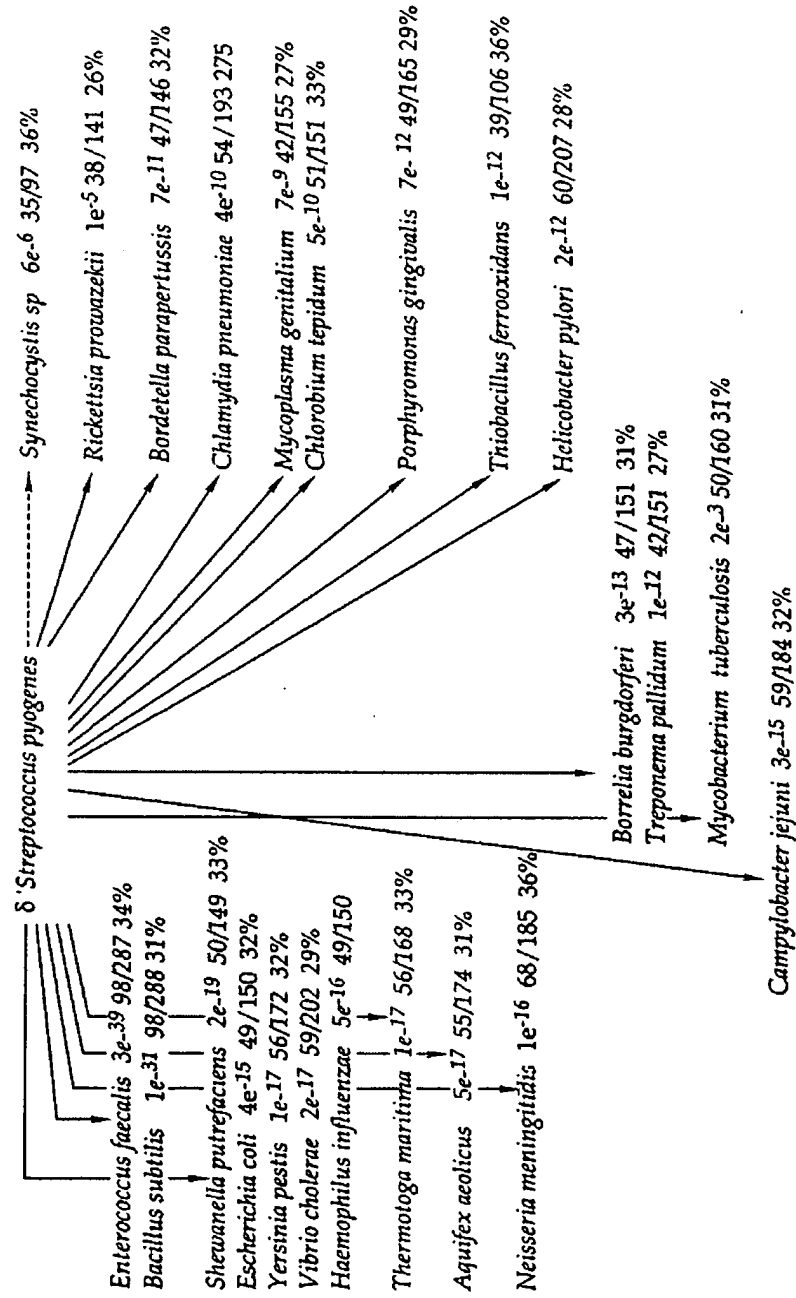


FIG. 20C

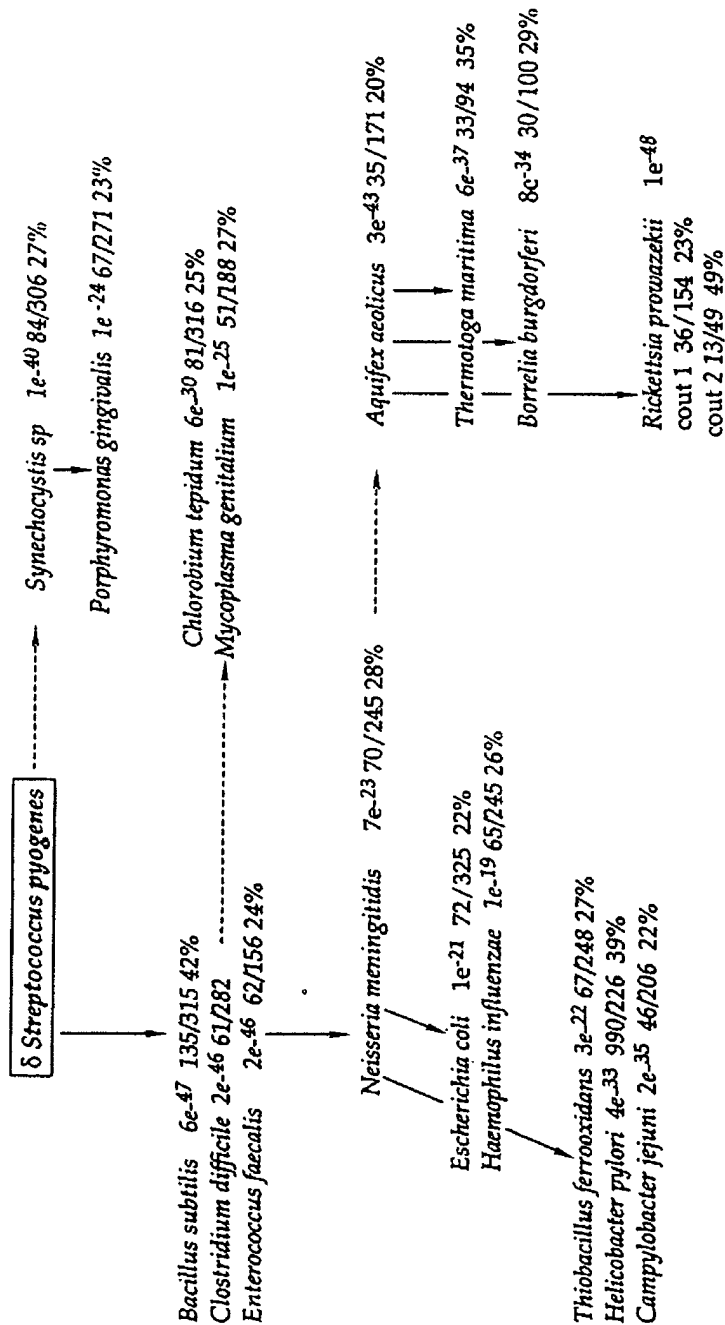


FIG. 20D

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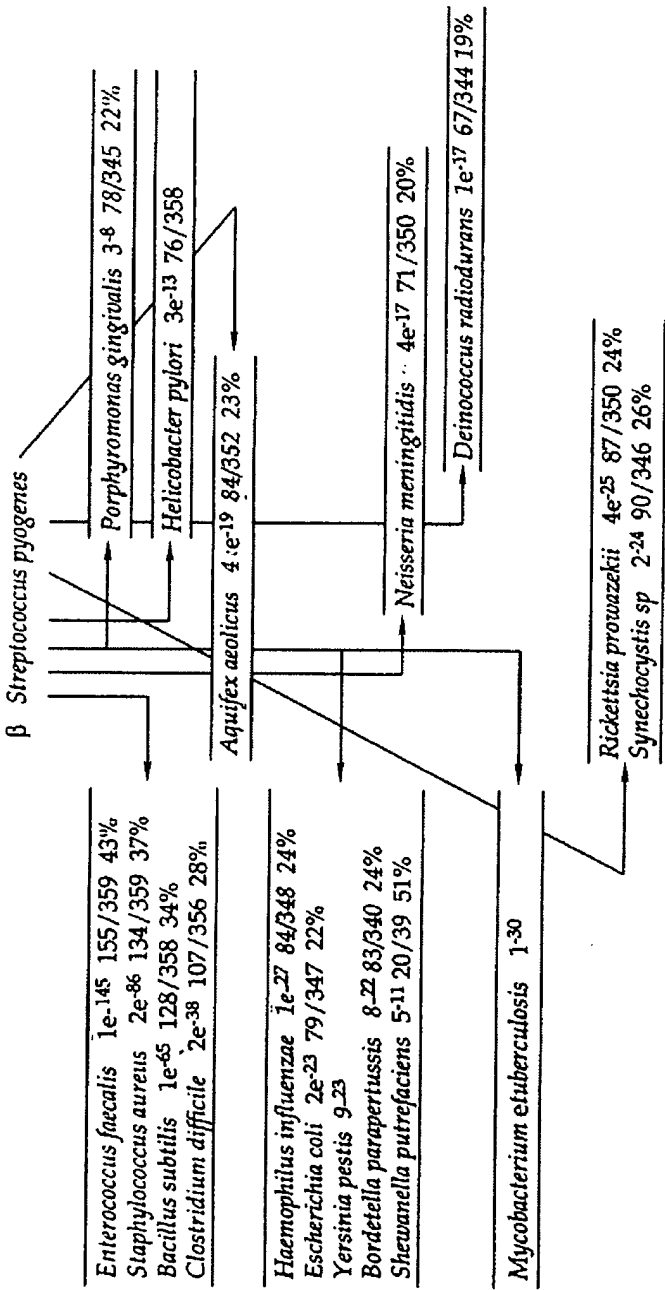


FIG. 20E

REPLACEMENT SHEET

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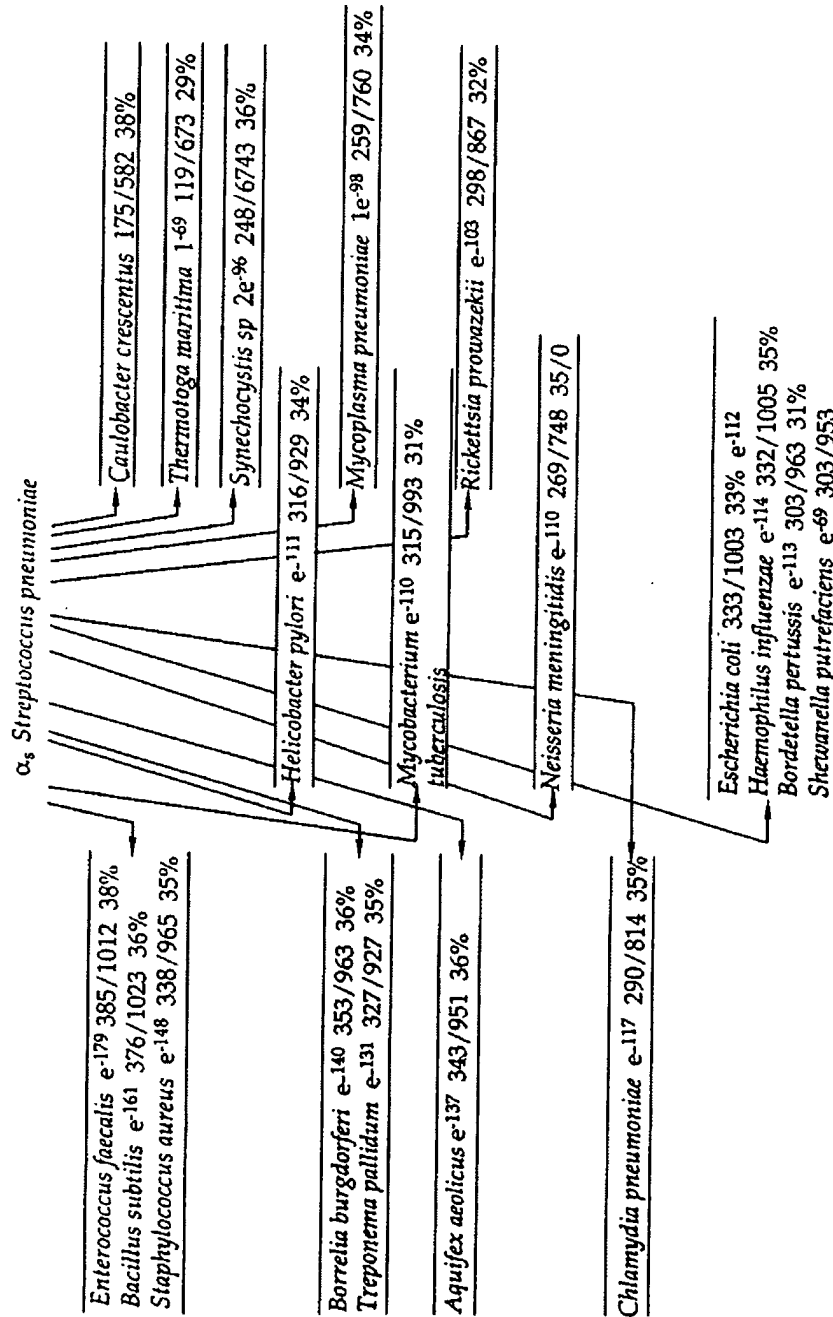


FIG. 20F

REPLACEMENT SHEET

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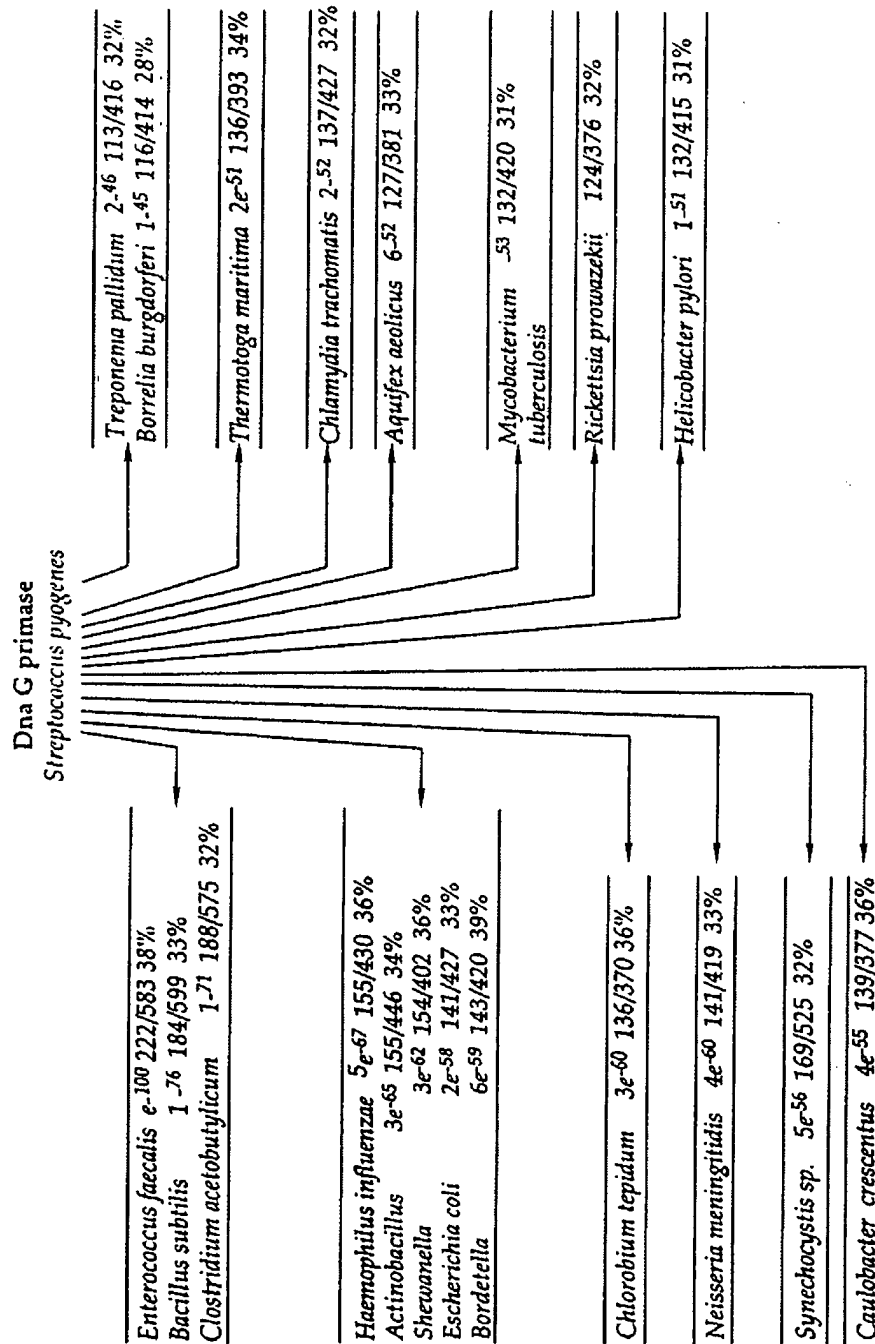


FIG. 20G

REPLACEMENT SHEET

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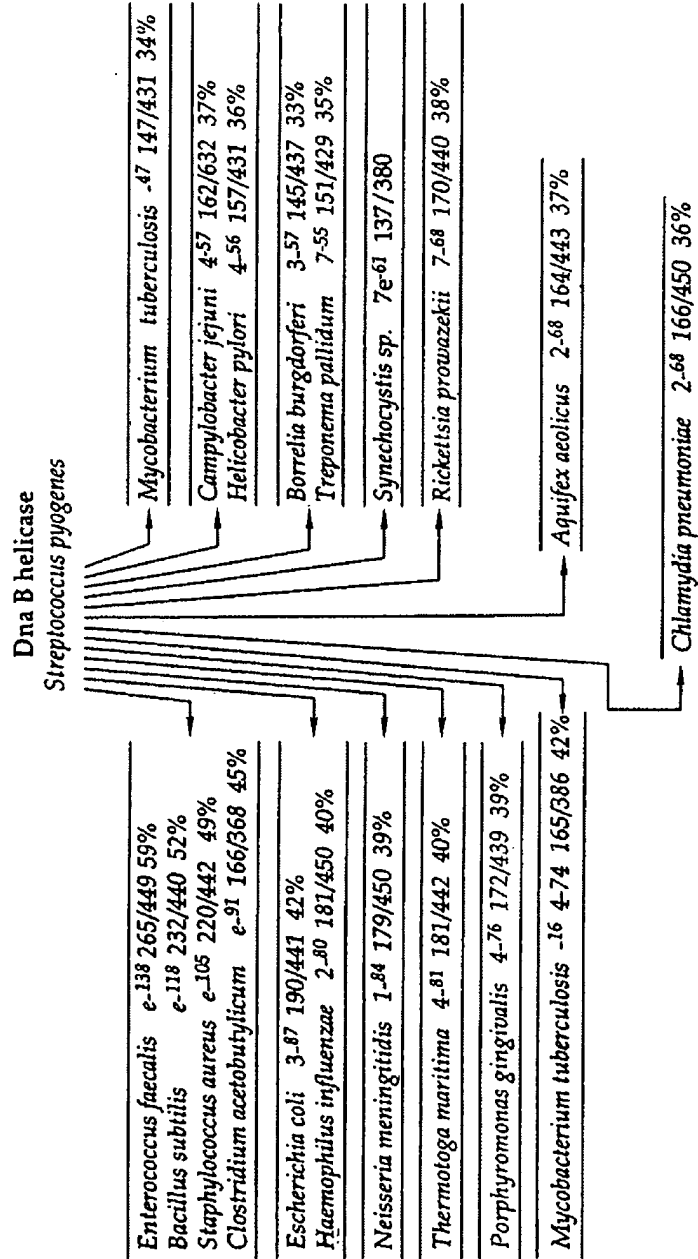


FIG. 20H

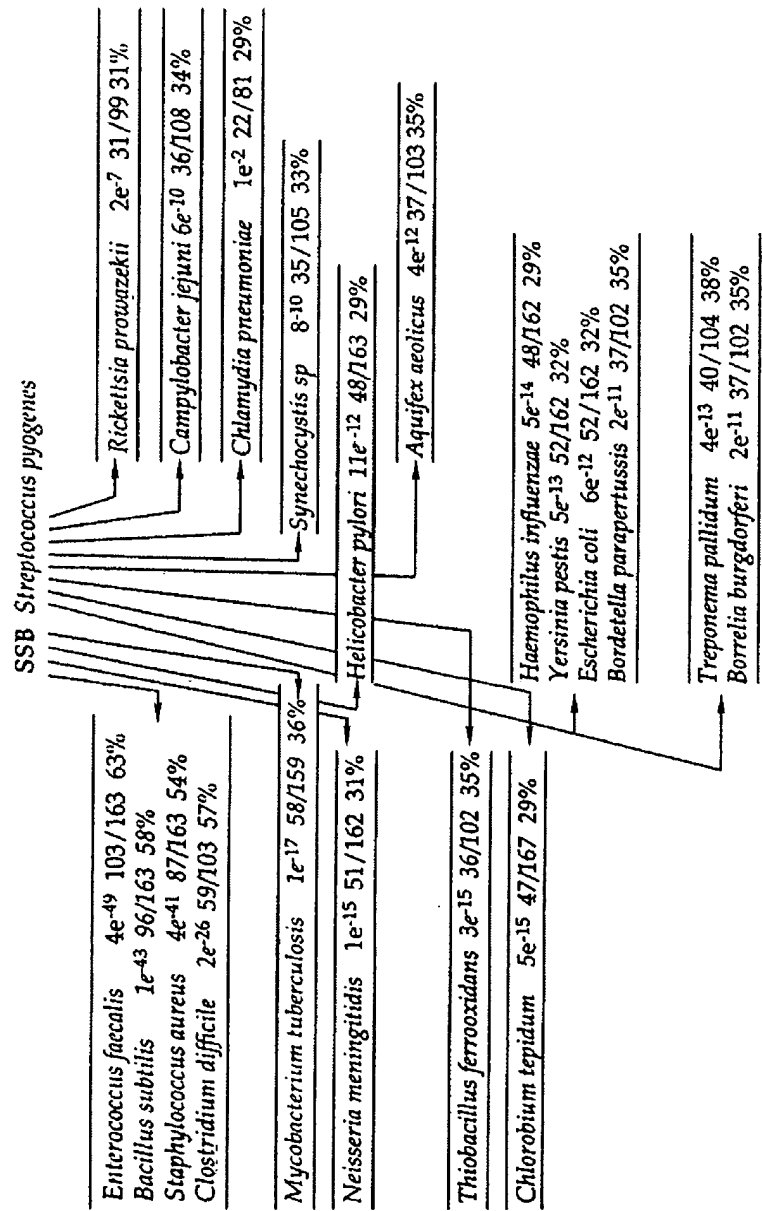


FIG. 201

FIG. 21A

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FIG. 21B

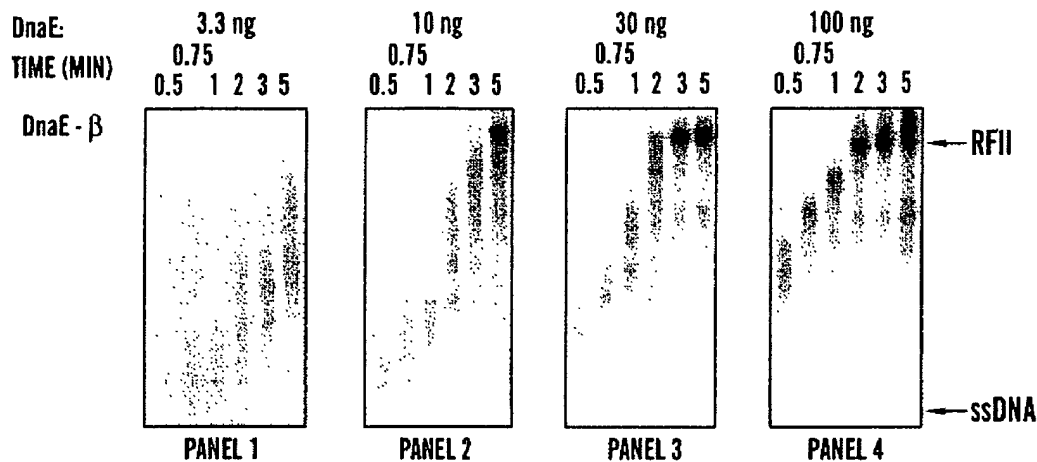
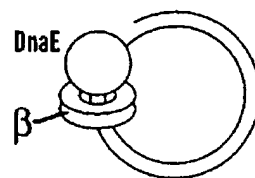
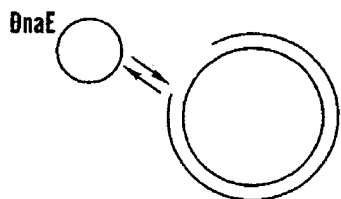


FIG. 21C

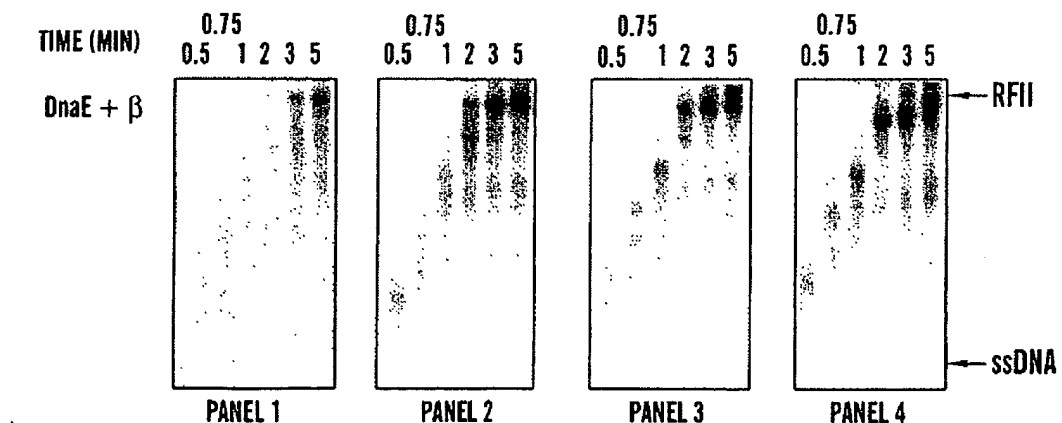


FIG. 21D

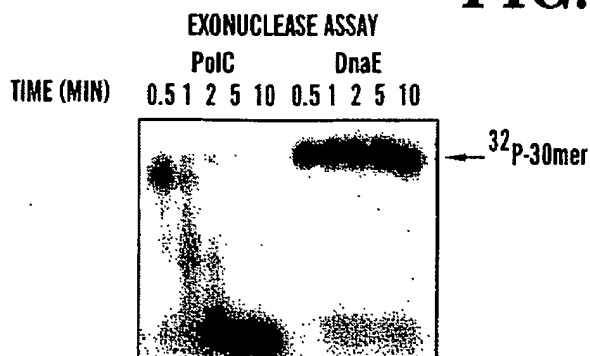


FIG. 21E

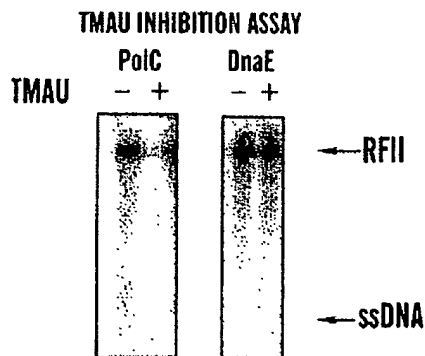


FIG. 21F